

Thu 18 May 2023 good_sequence_short.txt

Summary





Per tile sequence quality

Per sequence quality scores

Per base sequence content

Per sequence GC content

Per base N content

Sequence Length Distribution

Sequence Duplication Levels

Overrepresented sequences

Adapter Content

Basic Statistics

Measure Value

Filename good_sequence_short.txt

File type Conventional base calls

Encoding Illumina 1.5

Total Sequences 250000

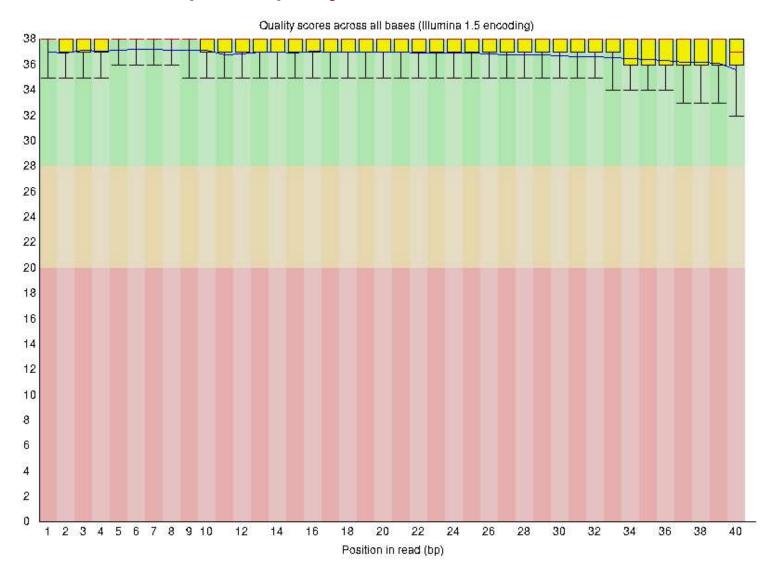
Total Bases 10 Mbp

Sequences flagged as poor quality 0

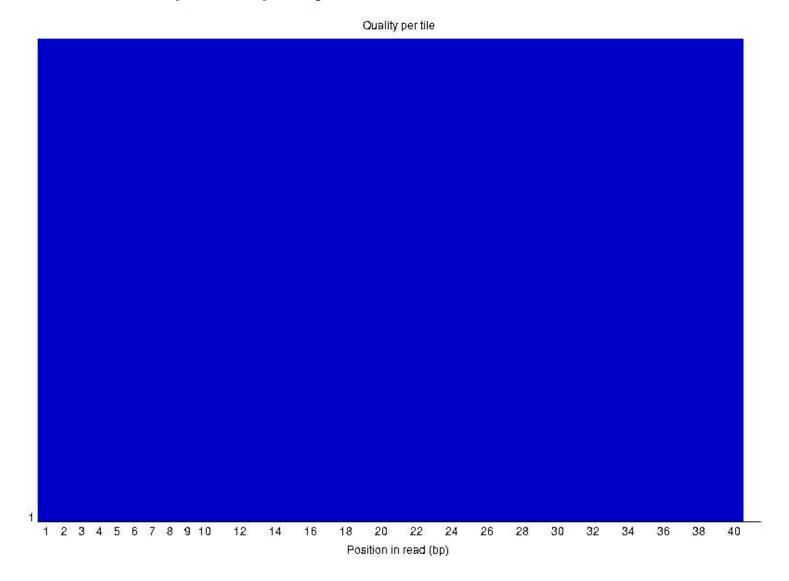
Sequence length 40

%GC 45

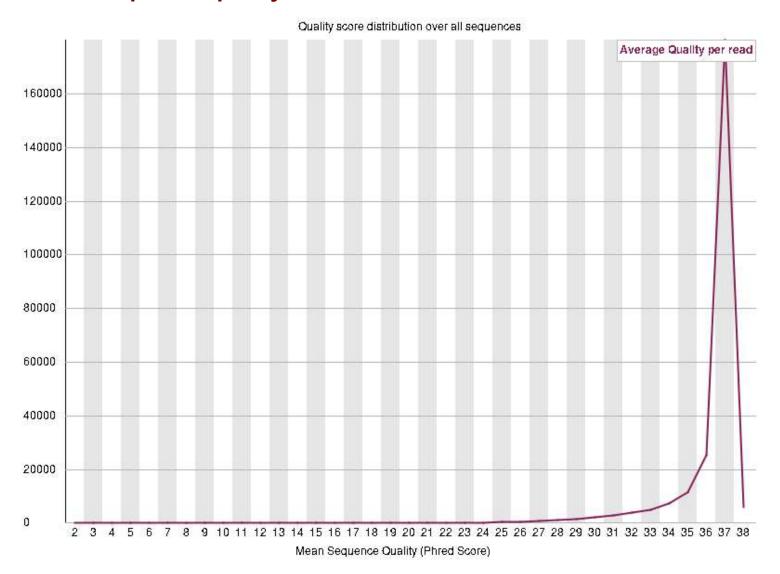
Per base sequence quality



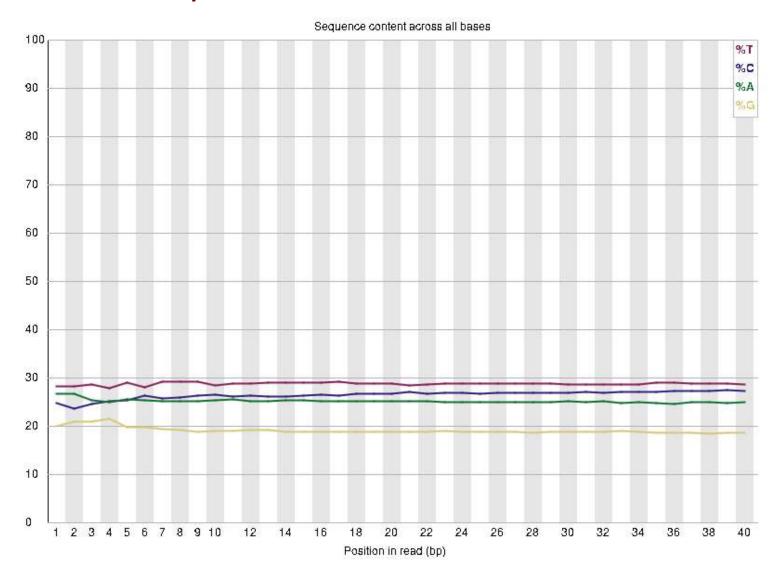
Per tile sequence quality



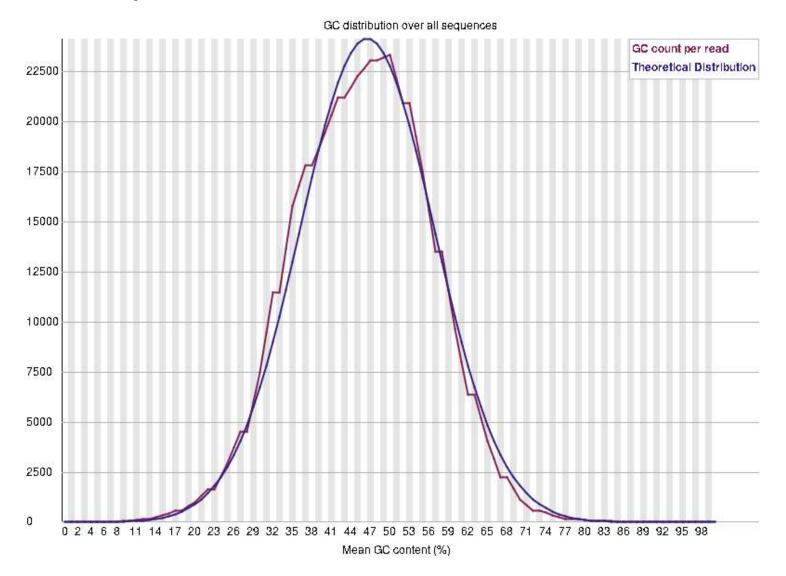
Per sequence quality scores



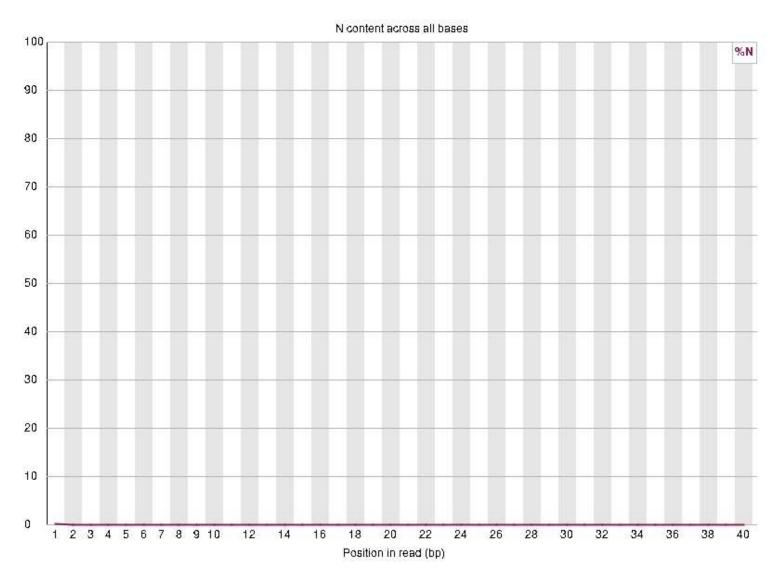
Per base sequence content



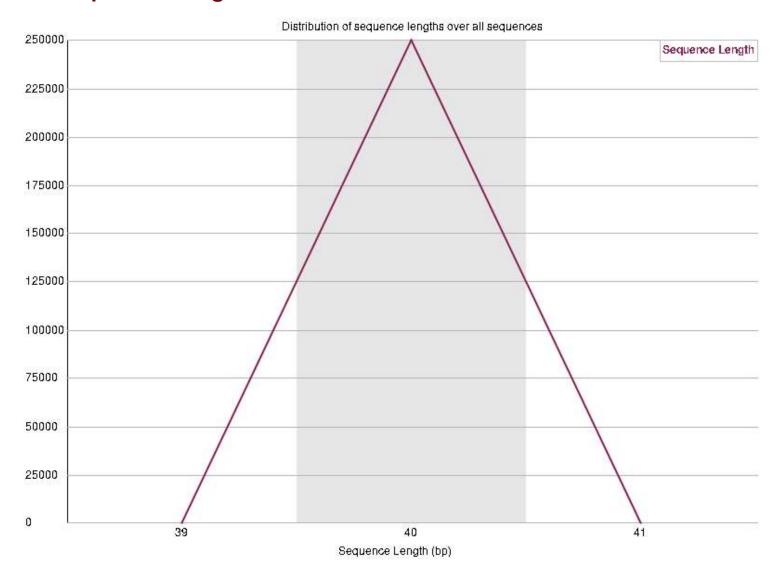
Per sequence GC content



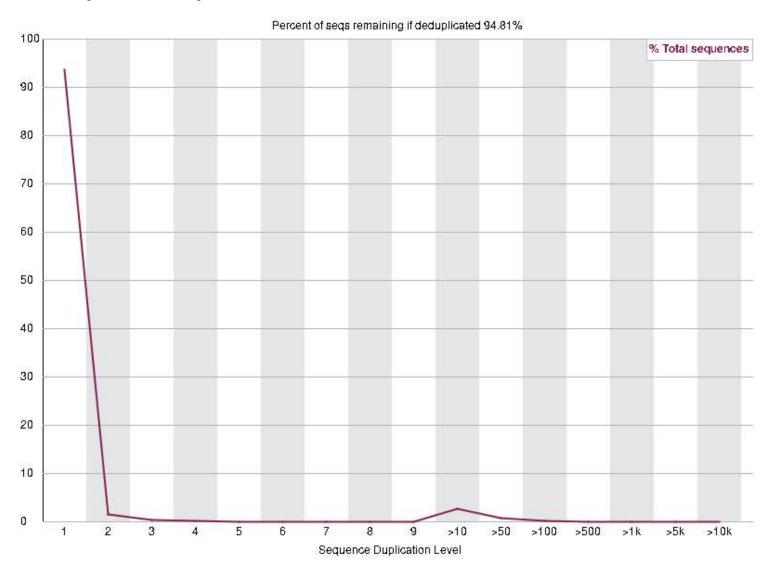
Per base N content



Sequence Length Distribution



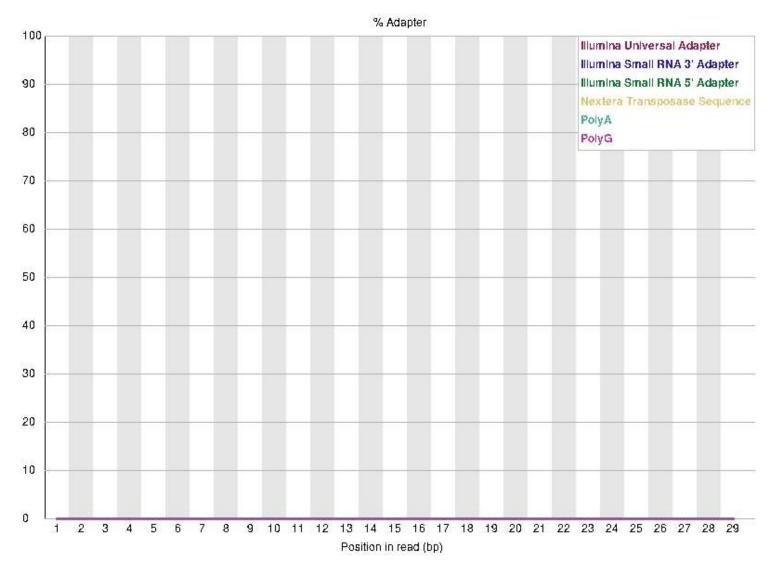
Sequence Duplication Levels





No overrepresented sequences

Adapter Content



Produced by FastQC (version 0.12.1)